

DDX-100US.ST25  
SEQUENCE LISTING

<110> Jonak, Zdenka L  
Trulli, Stephen H  
Tsui, Ping  
Lane, Pamela A.

<120> INTEGRIN LIGAND, HUMAN MINDIN

<130> DDX-100US

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<141> 2003-07-24

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<170> PatentIn version 3.2

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Gly Asn Trp Ser Gln Thr Ala Phe Pro Lys Gln Tyr Pro Leu Phe Arg  
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Pro Pro Ala Gln Trp Ser Ser Leu Leu Gly Ala Ala His Ser Ser Asp  
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Tyr Ser Met Trp Arg Lys Asn Gln Tyr Val Ser Asn Gly Leu Arg Asp  
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Phe Ala Glu Arg Gly Glu Ala Trp Ala Leu Ile Lys Glu Ile Glu Ala  
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Ala Gly Glu Ala Leu Gln Ser Val His Ala Val Phe Ser Ala Pro Ala  
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Trp Phe Val Gly Val Asp Ser Leu Asp Leu Cys Asp Gly Asp Arg Trp  
165 170 175

Arg Glu Gln Ala Ala Leu Asp Leu Tyr Pro Tyr Asp Ala Gly Thr Asp  
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Ser Gly Phe Thr Phe Ser Ser Pro Asn Phe Ala Thr Ile Pro Gln Asp  
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Thr Val Thr Glu Ile Thr Ser Ser Ser Pro Ser His Pro Ala Asn Ser  
210 215 220

Phe Tyr Tyr Pro Arg Leu Lys Ala Leu Pro Pro Ile Ala Arg Val Thr  
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## DDX-100US.ST25

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Glu Thr Pro Leu Asp Cys Glu Val Ser Leu Trp Ser Ser Trp Gly Leu  
 275 280 285

Cys Gly Gly His Cys Gly Arg Leu Gly Thr Lys Ser Arg Thr Arg Tyr  
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gcaggcggcc gagggcacag ggggtttcgc gctgctcctg accgcggtaa ggccgcgcgg	600
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20 25 30

Ser Ile Cys Ser Ala Arg Ala Leu Ala Lys Tyr Ser Ile Thr Phe Thr  
35 40 45

Gly Asn Trp Ser Gln Thr Ala Phe Pro Lys Gln Tyr Pro Leu Phe Arg  
50 55 60

Pro Pro Ala Gln Trp Ser Ser Leu Leu Gly Ala Ala His Ser Ser Asp  
65 70 75 80

Tyr Ser Met Trp Arg Lys Asn Gln Tyr Val Ser Asn Gly Leu Arg Asp  
85 90 95

Phe Ala Glu Arg Gly Glu Ala Trp Ala Leu Ile Lys Glu Ile Asp Gln  
100 105 110

Ser Pro Ala Ala Pro Gly Ala Pro Arg Ser His Gly Val Ser Gly Ala  
115 120 125

Pro Val Gln Ala His Ala Ala Gly Gly Arg Gly His Arg Gly Phe Arg  
130 135 140

Ala Ala Pro Asp Arg Gly Glu Ala Ala Pro Thr Ile Ser Ala Met Lys  
145 150 155 160

Gly Pro Leu Val Ala Gly Thr Gly Ile Gly Lys Gln Pro Pro Pro Phe  
165 170 175

Pro Thr Leu Leu Leu Arg Gly Pro Arg Val Pro Ser Ala Leu Ser Leu  
180 185 190

Leu Leu Leu Gln Asp Lys Val Ile Pro Lys Ala Pro Ala Thr Leu Asn  
195 200 205

Tyr Val Ser Leu Val Ile Ala Ala Pro Gly Asp Cys Pro Ser Ser Ser  
210 215 220

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Arg Gly Leu Ala Pro Thr Trp Leu Gln Ile Pro Gln Thr Trp Cys Ser  
225                    230                    235                    240

Arg Leu Cys Ala His Ser Pro Glu Gly Ala Ser Lys Arg Gly Pro Leu  
245                    250                    255

Glu Lys Ile Asn Gly Ala Val Ser Glu Ala Ser Val Phe Pro Cys Tyr  
260                    265                    270

Gly Ser Leu Cys Val Ile Lys Thr Ile Ser Val Ala Gln Lys Lys Lys  
275                    280                    285

Lys Lys  
290